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## Developmental Potential of Rat L6 Myoblasts in Vivo Following Injection into Regenerating Muscles

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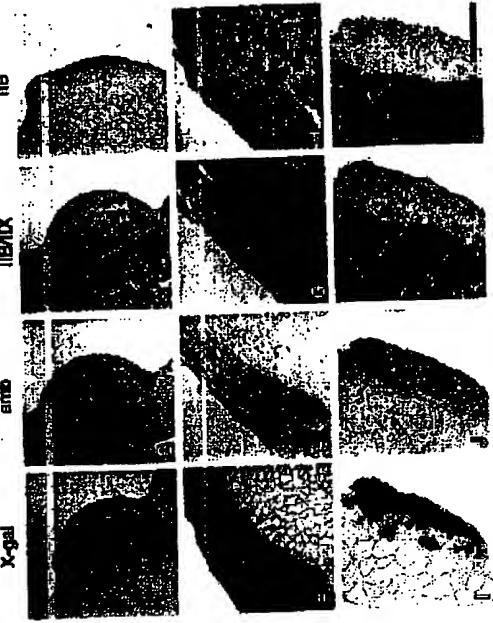
To examine the relative importance of envelopment lineage and environmental influences on the development of muscle fiber types *in vivo*, the phenotypy of muscle fibers formed from rat L6 myoblasts was examined following their injection into different regenerating adult muscles. Myoblasts were infected with a retrovirus carrying a LacZ reporter gene and their fate *in vivo* was examined using a panel of antibodies against various isoforms of myofibrillar MyHC isoforms. Since L6 myoblasts express LCK, following differentiation to cells, we wanted to determine if they would form IIX muscle fibers *in vivo* and whether their expression would alter *in situ*. Following injection, L6 cells either fused with each other or form homotypic fibers or fused with host muscle cells to form heterotypic fibers. Initially, bonemarrow fibroblasts expressed embryonic MyHC, while MyHC had replaced embryonic MyHC as the muscle transitioned to fast. We found that this transition was independent of donor L6 myoblasts (one host fast, IIA, and IIB fibers revealed that L6-derived muscle express embryonic and IIX MyHCs for up to 6 weeks postinjection, often as smaller domains surrounding L6 muscle). These results suggest that MyHC expression in muscle fibers derived from L6 myoblasts is regulated, in part, by intrinsic factors that limit the fiber type potential of these cells *in vivo*. © 1997 Academic Press

### INTRODUCTION

Adult mammalian muscle consists of several different fast fiber types (IIA, IIB, and IIX) and one slow after type (I), all of which can be characterized by differences in their speed of contraction (Schiaffino and Reggiani, 1996), resistance to fatigue (Gauthier, 1986), and pattern of myosin heavy chain (MyHC) expression (Armstrong and Phelps, 1984). In addition to the adult fast (IIA, IIB, IIX) and slow (type I) MyHC isoforms there are also several developmental isoforms, including embryonic and neonatal MyHCs, which are only expressed during muscle development (Condon et al., 1990; Hughes et al., 1990). The expression of the various MyHCs, including the down-regulation of the developmental iso-

forms, has been studied extensively in order to obtain insight into the mechanisms which regulate the development of the various muscle fiber types.  
Cross-innervation studies in which adult slow muscles were denervated and reinnervated by fast muscles demonstrated a transition in the muscle phenotype from slow to fast (Bulter et al., 1960). This suggested that the type of innervation received by the muscle governed the type of fiber phenotype. Experiments causing changes in the classical stimulation patterns of the muscle also indicated similar switches in MyHC expression (reviewed by Peres and Viborek, 1992), indicating that the changes caused by the nerve were due to the frequency of stimulation rather than specific trophic factors. Interestingly, cocultures of spinal cord and muscle produced the up-regulation of adult fast isoforms not normally present in cultured myotubes, even in the absence of synapse formation (Ercol-Prince et al., 1986). Therefore, direct innervation may not be necessary for the upregulation of the fast isoforms. These studies suggested that the environment regulates the pattern of MyHC expression in mature muscle fibers, with the nerve involved in the overall modification of that pattern. Recently, a growing body of evidence has suggested that

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**FIG. 1.** Pattern of myosin heavy chain (MyHC) expression in homotypic muscle fibers at 1, 4, and 8 weeks following infection of Lc myoblasts. A4 myoblasts were derived regeneratively from the phrenic muscle. Sections were either reacted with X-gel substrate (A, E, J) or analyzed for MyHC expression by immunohistochemistry with Myo 5B specific for embryonic (47A; B, F, I), late IIBMK (125; C, G, K) or late IIb MyHC (E, F, H, L). At 1 week postinfection (P1), X-gel staining reveals the presence of darkly staining, predominantly donor-derived myoblasts within the periphery of the muscle (A, J). These myoblasts react strongly for embryonic MyHC (47A; B, F, I) or other MyHC (C, G, K) (not shown) but do not react with the infected cells, suggesting that embryonic MyHC is the predominant isoform expressed in the homotypic fibers at this time. At 4 weeks P1 (G-E), most of the peripherally located MyHC is the predominant isoform expressed in the homotypic fibers at this time. At 8 weeks P1 (I-L), X-gel (C) MyHC but not IIb MyHC (H, L) nor for IIB MyHC (I), but not for IIB MyHC (J), reaction for embryonic MyHC is now the predominant isoform being expressed. The degree of reaction in homotypic muscle fibers was judged relative to the fiber-type specific reactions for IIBMK and IIB MyHC. In this study, Lc myoblasts infected with the Lc MyHC-A4 myoblasts, Bar = 370 nm in A-D, 270 nm in E-H, and 320 nm in I-L.

observed *in vitro* [DiMarzo *et al.*, 1983; DiMarzo and Stockdale, 1985]. However, these investigations did not determine if the homotypic myoblasts generated under these conditions became myoblasts. In addition, the developmental potential of these myoblasts could only be followed for 10 days, preventing the analysis of long-term environmental effects on this phenotype. In a different study, injection of C. C. C. myoblasts mouse satellite cells into the muscles of adult mice resulted in an alteration of the MyHC phenotype found *in vitro* when infected myoblasts fused with host muscle cells to form heterotypic fibers. The majority of MyHC isoforms found in cultures were down-regulated with differentiation of only one isoform typical of the muscle fibers into which these myoblasts were incorporated. These results supported the view that maturation ultimately controls muscle phenotype *in vivo* [Hughes and Blau, 1992]. Unfortunately, predominantly donor-derived fibers were not observed in this study and all of the labeled fibers appeared to be the result of the fusion of a small number of donor myoblasts with a substantially larger number of muscle fibers.

Since these two studies differed greatly in their design and in the type of fibers analyzed, it has not been possible to clearly define the relative contribution of intrinsic and extrinsic influences on the development of muscle fiber phenotypes *in vivo*. To address this problem, we have infected rat Lc myoblasts into regenerating hindlimb muscles of adult rats. The rationale for using Lc myoblasts is that this line expresses only two MyHCs, *in vitro*—embryonic fast and adult IIC MyHC—and may be committed to forming IIC muscle fibers *in vivo* [Wieser *et al.*, 1985; Pis and Merrifield, 1991]. The infection of these myoblasts with rat muscle tissue may establish patterns of MyHC expression prior to maturation and maturation.

To examine the effects of extrinsic factors on the phenotype of separate myoblast populations, investigations have been conducted characterized myoblast populations from various culture environments [Hughes and Blau, 1992; DiMarzo *et al.*, 1993]. Injection of quail embryonic myoblasts or satellite cells into developing chicken hindlimb muscle cells during differentiation [Blauchoff, 1986]. Consequently, the cells would have the option of fusing with each other or with muscle cells and fibers. Second, the derivation and

**Table 1**  
Myosin Heavy Chain Specificity of Monoclonal Antibodies<sup>a</sup>

Monoclonal antibody <sup>b</sup>	MyHC specificity	Isotype	Dilution	Indirect	Dilution ABC
47A <sup>c</sup>	Embryonic	IgG <sub>2a</sub> Rabbit polyclonal	1:10 1:50 1:100 1:13	1:10 1:200 1:14	1:10 1:350
NM62	Neonatal	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
M53.3 <sup>d</sup>	IIB	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
112P <sup>e</sup>	IIA	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
4A.7 <sup>f</sup>	IIA	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
SC-77 <sup>g</sup>	IIA	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
EM13 <sup>h</sup>	IIB	IgG <sub>1</sub>	1:50 1:53	1:50 1:5	1:50
EM32 <sup>i</sup>	All except emb. and IIA	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
SH2 <sup>j</sup>	Slow	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted
1D10 <sup>k</sup>	Slow	IgG <sub>1</sub>	Undiluted	Undiluted	Undiluted

<sup>a</sup> Antibodies obtained from (1) Pis and Merrifield (1991); (2) Butler-Browne *et al.* (1984); (3) Hughes *et al.* (1985b); (4) Sigma; (5) Donardi *et al.* (1993); (6) Boehringer *et al.* (1984). See text for details.

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muscle groups and their developmental potential was examined immunohistochemically over an 8-week period using a panel of antibodies specific for the different MyHC isoforms. Our results demonstrate that homotypic muscle fibers derived predominantly from Lc myoblasts maintain their *in vitro* pattern of MyHC expression since they maintain detectable levels of only embryonic and IIC MyHC. Interestingly there is a transition in the phenotype of these muscle fibers as they develop. The innervation of such myofibers, starting at 4 weeks postinfection, has previously been observed in experiments where large-scale degeneration occurred prior to myogenesis when limb transplantation [Pis and Merrifield, 1991]. The innervation of individual fibers can be assessed by examining the status of individual fibers can be assessed by examining the expression of neural cell adhesion molecules (NCAM) along the surface of the fiber, since NCAM is localized along the entire length of intact muscle fibers but becomes localized exclusively to the motor endplate following innervation [Covault and Sanes, 1985; Covault *et al.*, 1986].

In this study, Lc myoblasts infected with a constitutively expressed Lc-Z reporter gene were injected into different

**TABLE 2**  
Myotin Heavy Chain expression of Heterotypic L6AG-A4 Myotubes

Weeks after infection	Muscle	Myotin heavy chain expression					
		Embryonic	Neonatal	DIA	IM	IX	Show
1	Gastro.	+++	-	-	-	-	-
	Soleus	+++	-	-	-	-	-
	Plantaris	+++	-	-	-	-	-
	TA	+++	-	-	-	-	-
2	Gastrocn.	+++	-	-	-	-	-
	Soleus	+++	-	-	-	-	-
	Plantaris	+++	-	-	-	-	-
	TA	+++	-	-	-	-	-
4	Gastrocn.	++	-	n.d.	n.d.	n.d.	n.d.
	Soleus	++	-	n.d.	n.d.	n.d.	n.d.
	Plantaris	++	-	n.d.	n.d.	n.d.	n.d.
	TA	++	-	n.d.	n.d.	n.d.	n.d.
8	Gastrocn.	++	-	n.d.	n.d.	n.d.	n.d.
	Soleus	++	-	n.d.	n.d.	n.d.	n.d.
	Plantaris	++	-	n.d.	n.d.	n.d.	n.d.
	TA	++	-	n.d.	n.d.	n.d.	n.d.

Note: +++, >90% of myotubes are positive; ++, <90% of myotubes are positive; +, <1% of myotubes are positive; n.d., not determined.

mixture of different type II fibers) (Armstrong and Phillips, 1984), the expression of L6 MyHC was maintained, often in conjunction with the MyHC isoforms characteristic of the fiber type. However, in heterotypic fibers resulting from the incorporation of L6 myoblasts mixed into slow type I fibers, L6 MyHC was only transiently expressed. These results suggest that MyHC expression in muscle fibers formed from L6 myoblasts is regulated, in part, by intrinsic factors that limit the fiber type potential of these cells *in vivo*.

## MATERIALS AND METHODS

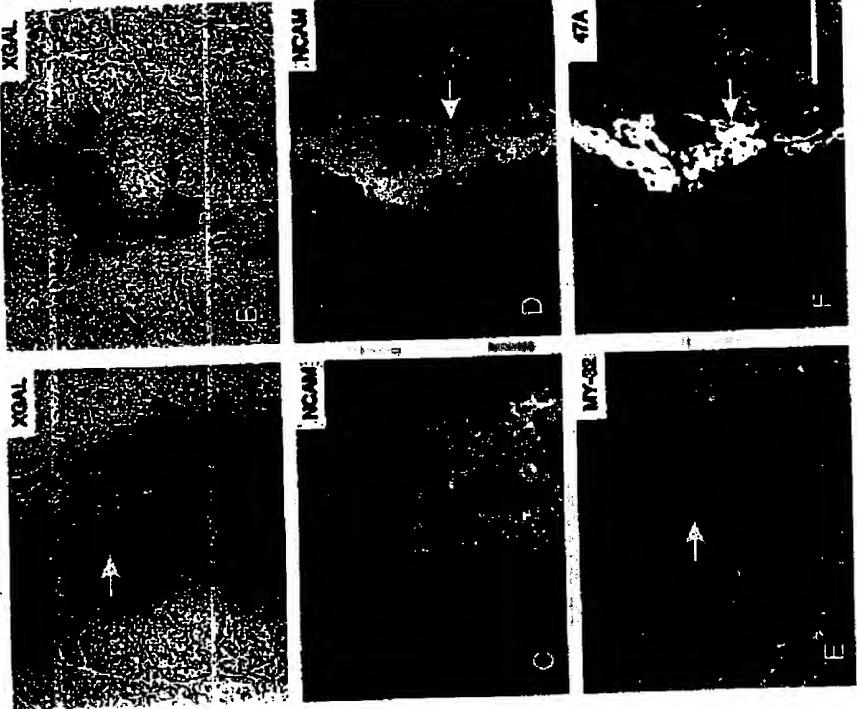
### Infection of L6 Rat Myoblasts with a β-Galactosidase Reporter Construct

A subclone of the L6 rat myoblast cell line originally isolated by Yaffe (1961) was obtained from Dr. B. D. Samuel (Department of Biochemistry, University of Western Ontario) and grown in complete DMEM containing 10% fetal calf serum. Cells were tested for the ability to take up [3H]thymidine, and duplicate wells were tested for β-galactosidase (β-gal) activity. Clones to which all cells expressed high levels of β-gal before and after fusion were tested for regenerating adult rat muscles.

### Infection of L6 Myoblasts into Regenerating Adult Muscles

Once stable clones of L6 myoblasts were obtained in which a high constitutive level of β-gal could be observed, cells were expanded to obtain large populations for injection. One each clone, grown for 1 day (approximately 10<sup>6</sup> cells/100 mm dish), was rinsed twice with DMEM medium and harvested for another 4 h. Conditioned medium containing the L6C reporter was harvested, centrifuged at 800 g for 10 min, and filtered through a 0.45-μm filter to remove cells and assayed in 10-ml sterile aliquots at -80°C.

To infect 16-month-old mice, low passage cells (11P7775; Rock)



**FIG. 2.** Immunofluorescent localization of NCAM and MyHC expression in humotrophic myotubes 7 days after injection of L6AG-A4 myoblasts into the extensor digitorum longus muscle of adult rats. Serial sections were either stained for X-Gal (A, B) or analyzed using immunofluorescence with an NCAM-specific polyclonal antibody (C, D) or a MyHC-specific monoclonal (I) or a MyHC-specific (F, G) secondary antibody (F, G). Primary antibodies were identified by fluorescein isothiocyanate (NCAM) and FITC (MyHC). All are all NCAM positive (A, B) and MyHC expression (I) but not near (D, E) MyHC at this time. Bar, 55 μm.

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Hanks Balanced Salt Solution (CBFS), then rehydrated with 1 ml 10 dilution of 2.5% cryoin CMB-FBS until all of the cells lifted off the plate. The harvested cells were then collected and washed twice with CMB-FBS. The resulting pellet was resuspended in a medium containing 0.1% penicillin/streptomycin, 0.1% amphotericin B, 0.25 mg/ml trypsin inhibitor, 10 mM EGTA, 10 mM pyruvate, 10 mM sodium pyruvate, 100 U/ml penicillin, 10 mM NaCl, and 0.1 M sodium phosphate, pH 7.5, for 10 to 37°C, as described by Sharnabha et al. (1989). Clones were isolated using cloning cylinders, expanded, and duplicate wells were tested for high levels of β-gal before and after fusion were tested for their ability to take up [<sup>3</sup>H]thymidine.

### Injection of L6 Myoblasts into Regenerating Adult Muscles

Once stable clones of L6 myoblasts were obtained in which a high constitutive level of β-gal could be observed, cells were expanded to obtain large populations for injection. One each clone, grown for 1 day (approximately 10<sup>6</sup> cells/100 mm dish), was rinsed twice with DMEM medium and harvested for another 4 h. Conditioned medium containing the L6C reporter was harvested, centrifuged at 800 g for 10 min, and filtered through a 0.45-μm filter to remove cells and assayed in 10-ml sterile aliquots at -80°C.

To produce transfecting L6C virus, PH1.2 L6C cells were grown for 1 day (approximately 10<sup>6</sup> cells/100 mm dish), then collected and washed twice with DMEM medium, rinsed once with Ca<sup>2+</sup>, Mg<sup>2+</sup> free

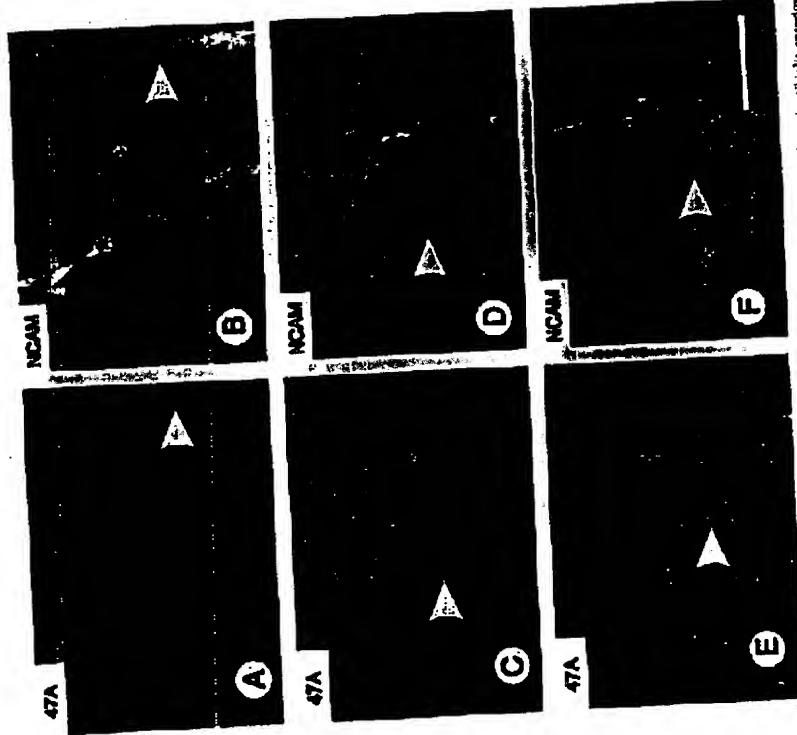
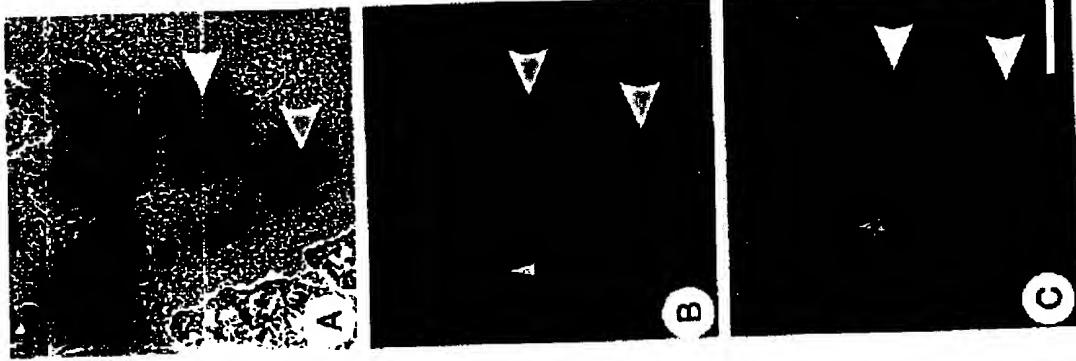


FIG. 5. NCAM expression in hemispheric fibers 50 days after infection or in cultures maintained against embryonic MytG (47A, A, L-1) and adult NCAM (B, B). Primary antibodies were carried out with antibodies conjugated to biotinylated NCAM (NC-10) or rhodamine-conjugated secondary antibodies. The stain for embryonic MytG can be seen in A and B, while myelin basic protein stains are located in C and D. The lack of condensation between NCAM and NCAb can be seen in C and D, and E and F. All myelin basic protein stains are located in G and H. The downregulation of embryonic MytG and NCAM strongly suggests that myelinium is not involved in the expression of embryonic MytG and NCAM.

tion of medium periorbital and cerebral lesions. Their hindlimbs were then shaved and washed with alcohol prior to injection. Approximately one million cells in 50 µl of the cocktail were injected into each of three sites—(a) the sciatic and gluteal, (b) the gastrocnemius, and (c) the orbicularis anterior and extensor digitorum longus muscles of the right leg. Control infections, consisting of

Chloralose (chloral hydrate) was administered orally but without effect. A 1-ml syringe with a 16-gauge needle was inserted on the left ear. A 1-ml syringe with a 16-gauge needle was used to perform the injections. The animals were allowed to recover under heat lamps and then maintained as an isolated colony. Each animal received a daily injection of cytoporph. A Sanjour (Curaids Inc., Montreal, Quebec) concentration of 15 mg/kg parenteral

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**FIG. 4.** Coexpression of MyfIC and NCAM expression in bovine myoblasts 12 days after infection into the white gastronemius muscle of adult Bov. Serial sections were analyzed for  $\beta$ -galactosidase staining (blue) and for MyfIC/NCAM coexpression using immunoperoxidase reaction. Colocalization (B, C, NCAM was detected with a rabbit polyclonal antibody using a fluorescein-conjugated secondary antibody, and MyfIC was detected with a mouse monoclonal antibody and a biotin-conjugated secondary antibody. Avidin-biotin-peroxidase complex was visualized with diaminobenzidine. Colocalization reveals two myofibers (arrowheads) in a bovine muscle where MyfIC is localized in the nucleus of the muscle fiber, but not in the cytoplasm. The arrows point to two other myofibers where NCAM with MyfIC colocalizes in the cytoplasm. The absence of MyfIC reactivity in these myofibers indicates that they are coexpressing MyfIC with NCAM. This suggests that the differentiation is not essential for the expression of MyfIC. MyfIC is also expressed in the somite muscle tissue. The homogenate prepared from L6E9G4M-derived homotypic fibroblasts (the blue dots) show no MyfIC staining. The upper left of each panel (A) stains with MyfIC antibody (the red dots). The right panel (C) indicates that it expresses endogenously MyfIC. Since this is negative for NCAM, this is an example of a differentiated heterotopic fibroblast which continues to express MyfIC. MyfIC: 50  $\mu$ m.

## **Immunohistochemical Analysis of MyHC Expression in Muscle Fibers Containing Dendro-L6 Nuclei**

body weight. After 4 weeks, cyclophosphamide injections were reduced to 7 days to reduce cramps on the tail. Six rats were sacrificed at 7, 14, and 28 days, 4 rats were sacrificed at 40 and 56 days, and one rat was sacrificed at 61 days. At day 61, rats were anesthetized with isoflurane gas and killed by exsanguination. Superior vena cava was cannulated with 18 gauge silicon tubes. Superior vena cava was identified by finding the vein that drains in the right atrium. The catheter was embedded in Tissue Tek OCT freezing compound and stored sectioned at 10-15  $\mu$ m on a Leitz cryotome. Every tenth section was placed in 1% glutaraldehyde in PBS, and embedded in Durcupan resin. A total of 10-15 pairs of tissue sections were taken from each rat. Tissue sections were stained with 1% uranyl acetate and lead citrate for electron microscopy.

Solutions were examined for MyHC expression using protein kinase C inhibitors [A,B,C] fluorescence or AOC-alkaline phosphatase immunohistochemistry. Sections were blotted at 10% graft section in phosphate buffered saline (PBS) for 30 min at 37°C, incubated in primary antibody (for Ig G or IgM) washed with PBS. Primary monoclonal antibodies 47A, IgM S32, 212R, IgG 47c, SC-71, IgG 8E, IgG 8F, and IgD100 were used for these analyses. In addition, rabbit polyclonal antibody N56 was used, which specifically recognizes neuronal MyHC (Hunter and Whalen, 1984). The specificity and optimal dilutions of these antibodies is summarized in Table 1. Sections were then incubated with a secondary antibody (1:10,000 dilution of either biotinylated goat anti-rabbit IgG (HRP conjugate), IgG 47A, streptavidin) or rabbit anti-goat IgG (HRP conjugate) for 1 hr at 37°C (ICAR DGC, Tago Inc., Burlingame, CA). Biotinylated streptavidin (Vectastain Elite kit, LS Plus, Elite kit, Vectastain, Burlingame, CA) was used to detect the biotinylated antibodies.

**Immunofluorescent Colocalization of MyHC Isoforms and NCAM**

Once injection sites containing doxorubicin were identified based on *S. gal* immunohistochemistry, serial sections were analyzed for the presence of MyHC isoforms.

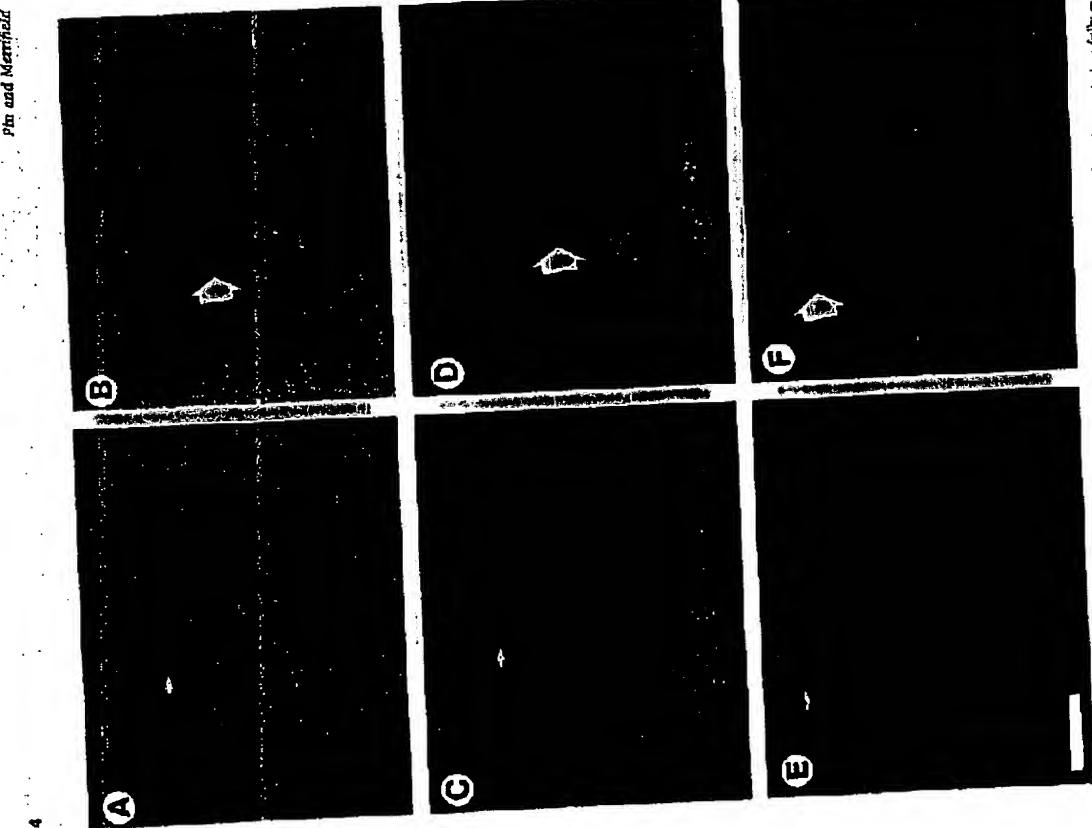


FIG. 5. Rat diaphragm muscle fibers viewed at low [A, C, E] or high [B, D, F] magnifications. Serial sections were characterized for X-gal staining [A].

tried for (a) the MyHC phenotype of fibers containing donor nuclei, and (b) the presence of NCAM molecules along the surface of the fibers. These sections were fixed with 3% methanol for 6 min at 37°C, rinsed briefly in PBS, then stained with 1:25 goat serum to block non-specific antibodies and the secondary antibodies. Subsequent incubation of the primary antibodies and the secondary antibodies used in their detection was carried out. Slices were first incubated for 1 hr at RT in IgG, antibodies (1:4 and 1:8) followed by several rinses of PBS and 1-hr incubation in BT with fluorescein-conjugated goat anti-rabbit IgG, (NCN Biomedicals Canada Ltd., Montréal, Québec, diluted to 1/50 in PBS containing 0.1% bovine serum albumin (BSA). After several rinses with PBS, slides were incubated for IgG antibodies (IgG 1:4 and IgG 1:8). These antibodies were detected using a rhodamine-conjugated sheep anti-sheep (SAM) IgG, (diluted to 1/50 in PBS-SAM). Slices were counterstained with a 5% glacial solution in PBS containing 5% propidium iodide and 0.5% Hoechst dye.

To characterize the innervation status of fibers containing donor nuclei, a rabbit polyclonal antibody that recognizes all forms of NCAM (kindly provided by Dr. Gérardine Bouet, CNRS, Marcellis, France; Bouet and Marchal, 1986) was used in combination with the various mouse monoclonal antibodies. Since the NCAM antibody is a rabbit polyclonal antibody and the MyHC-specific antibodies are mouse monoclonal antibodies, both primary antibodies were treated simultaneously following incubation in the primary antibodies for 1 hr at RT, sections were rinsed several times with PBS and incubated in a 1:30 dilution of both fluorescein (FITC)-conjugated goat anti-rabbit and rhodamine (RITC)-conjugated anti-monoclonal (RAM) L6C secondary antibodies (NCN Biomedicals Canada Ltd., Montréal, Québec, Ontario) in PBS-BSA for 1 hr at RT. Sections were rinsed several times with PBS and then coverslipped as described above.

#### Determination of Fiber Types Based on Myosin Heavy Chain Expression

Following ABC-AP immunolocalizations using MyHC-specific Mabs, sections were analyzed for the number of host/donor heterotypic fibers (which stained for X-gal and showed characteristic cytosolic or nuclear fibers) and host-derived mature fibers. To ensure accurate counting, only fibers clearly belonging to one group or another were scored. Scoring of host-derived fibers was limited to the area immediately adjacent to the injection site so that regional differences in the muscle would be minimized. To provide accurate counts, specific muscle fascicles were analyzed in each serial section. The number of positive fibers within these areas was determined after labeling with 1:120 (MBP/DC), 1:174 (MAB, SC-71 (MAB) 1:25 (all MyHCs except embryonic and DC), 1:52 (IBA), and 1:10 (D10) Mabs. Fibers were then classified as types I, IIa, IIa/IIx, IIx, IIb/IIx, or IIb. To determine the number of fibers belonging to each group the following equations were used: [A] Total = No. of X-gal positive fibers [B] (above) = No. of 100% positive fibers [C]

[A] Total No. of fibers [B] (D)/[A]X - No. of 100%-positive fibers - No. of Type Ia fibers [C] (D)/[A]X - No. of 100%-positive fibers - No. of IB-EQ fibers. Since No. f. Ia, IIa, fibers = 100% fibers, since no 100% fibers can be counted both IB and IIa, it was impossible to obtain an accurate determination of the number of 100% fibers. However, Mab 1A-74 has been shown to correlate well with IB MyHC and higher levels of expression. Therefore, it was possible to obtain a lower limit to the number of fibers containing both IB and IB MyHC using the equation:

$$\text{IB}/\text{IBX} = \text{No. of 4A, 74-positive fibers} - \text{No. of IIa fibers} - \text{No. of IB/IX fibers} - \text{No. of IIa/IX fibers}.$$

Therefore, the number of IB fibers could be determined by:

$$\text{III} = \text{No. of 100%-positive fibers} - \text{No. of IB/IX fibers}.$$

From these equations the percentage of fibers belonging to each class was determined for regions of the muscle within and outside of the injection sites.

#### Photography

All images were captured using a Zeiss microscope and the computer software program Numbered Images. Pictures were produced using a Phaser 440 Tektronix dye sublimation printer.

#### RESULTS

#### Characterization of L6AG-4A-Derived Heterotypic Fibers after Injection Into Regenerating Adult Muscles

To study the effects of various environments on the maintenance of the L6 phenotype, several different muscles were targeted for injection. The lateral (white) portion of the gluteus communis is composed predominantly of IB fibers. The iliobial tensor and plantaris muscles represent typically adult muscles containing all adult fiber types while the rodent muscles of the gastrocnemius and the soleus muscle represent typically slow muscles (Armstrong and Pheifer, 1984). These last two muscles contain large numbers of slow and fast fibers and small numbers of IB fibers and therefore represent muscle environments in which maintenance of the L6 phenotype may be most challenged.

Following injection, L6 myoblasts formed both heterotypic and bimimetic fibers. Heterotypic fibers result from the fusion of donor cells with each other to form new myo-

blasts or MyHC expression using ABC-fluorescence using MyHC Mabs 47/A, C and D) or monoclonal MyHC (MABs 1A-74, SC-71 (MAB), and F1, and primary antibodies were identified with a rhodamine-conjugated secondary antibody. X-gal staining is fully uniform (A and B) and shows donor cell fusion to two adjacent fibers (C). One of the fibers maintains some degree of embryonic MyHC expression localized only to the periphery of the fiber (C and D), with some lighter staining throughout the rest of the fiber. The absence of normal MyHC (E and F), along with peripheral nucleation revealed by Hemacryl dye staining (E) suggests that this fiber is fully enucleated and under normal circumstances would not express embryonic MyHC. Bar, 100 μm (A, C, E) or 40 μm (B, D, F).

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## **Characterization of NCAM Expression in Fibroblasts and its Relationship to Synaptic Fibers and its Relationship**

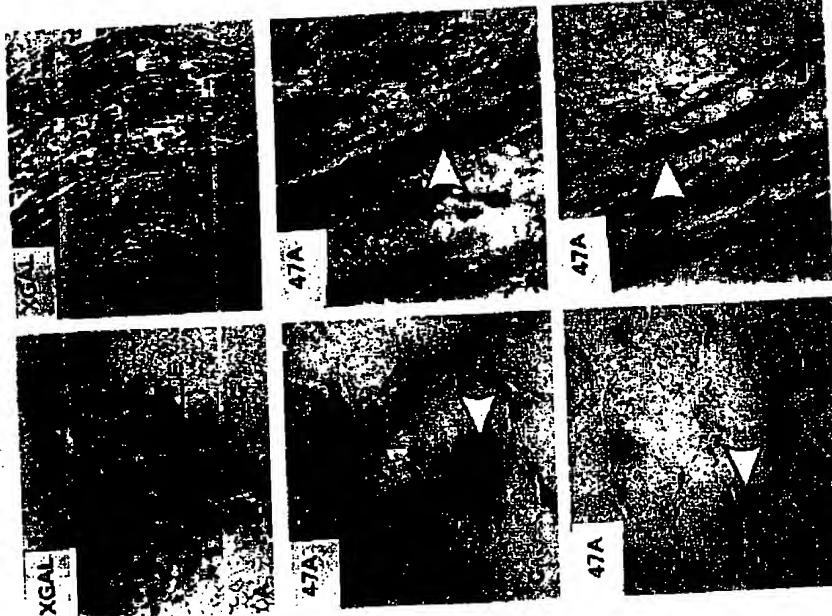
**to MyHC Expression**

Because the loss of embryonic MyHC in some heart tissue was observed in all insertion sites, we wanted to examine the role of myosin in this transition. To address this issue, a polyclonal rabbit antibody, specific for all NCAM isoforms, was utilized along with the various MyHC antibodies. NCAM is known to be expressed along the surface of cardiocytes prior to lamination. Upon inner-

tion, NCAW becomes localized in the epineurium [Figures 3a and 3b, 1922; Govaris and Sánchez, 1985]. Therefore, myeloses that are NCAW negative are likely observed while those that show punctate staining along the membrane are not immunoreacted.

X-gal immunohistochemistry revealed high levels of  $\beta$ -gal and had a circular cross-section, typical of donor-derived myotubes. Immunofluorescent localization of serial sections using an NCAM-specific polyclonal antibody and a monoclonal antibody specific for embryonic MyoTc revealed that NCAM and embryonic MyoTc were specifically coexpressed in these cells. This suggested that donor-derived myotubes were not in an early time after infection. Contrariwise, the two antibodies to muscle fibers outside the infection site indicated that the host fibers were undergoing regeneration, a process that can involve both dedifferentiation and reexpression of developmental programs. Fluorescent immunohistochemistry revealed high levels of  $\beta$ -gal and had a circular cross-section, typical of donor-derived myotubes. Immunofluorescent localization of serial sections using an NCAM-specific polyclonal antibody and a monoclonal antibody specific for embryonic MyoTc revealed that NCAM and embryonic MyoTc were specifically coexpressed in these cells.

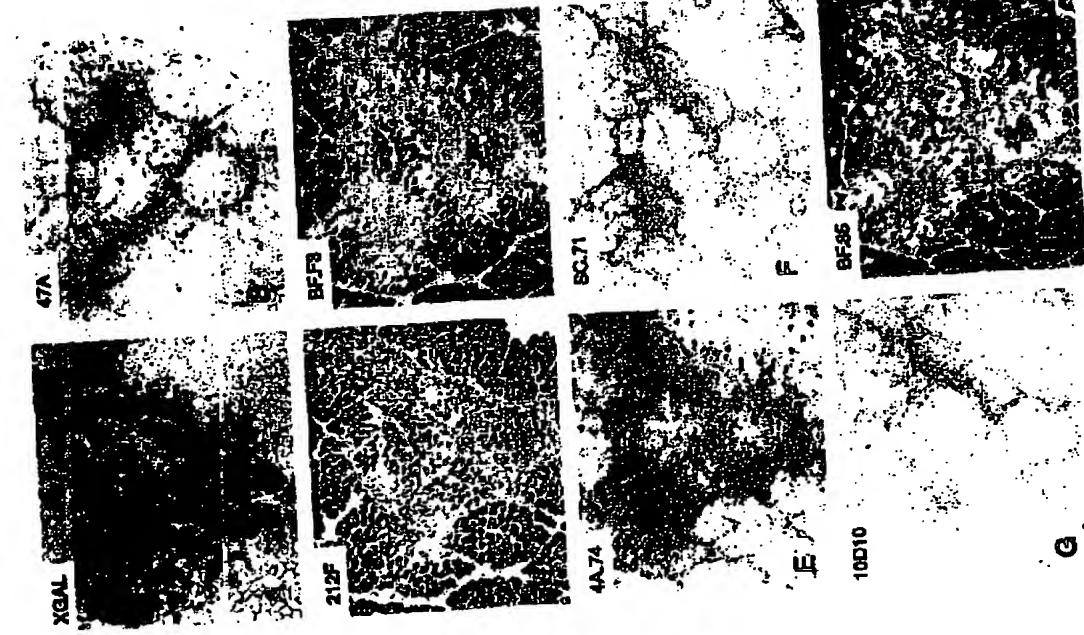
Examination of Impression



**FIG. 6.** Nuclear laminae of embryogenetic Myotis persicus at adult size after injection of  $\text{D}_2\text{O}$  (top) and  $\text{H}_2\text{O}$  (bottom). Higher X-ray luminescence intensity revealed large areas of heterochromatin fibers in the nucleus of embryogenetic Myotis within adult size epiglottides of the larynx under air (C, E) and  $\text{D}_2\text{O}$  (F) 50 days after injection. These domains surrounded single nuclei (G) and ranged between fiber fibers in the nucleus under air (H, I) and  $\text{D}_2\text{O}$  (J, K) 50 days after injection. These fibers also reacted with M7-32 IgG (not shown), indicating that they have an adult fat phenotype. Heterochromatin fibers become the larger fibers can also react with [-] Bar, 500 nm for A and B, 40 nm for C and E, and 80 nm for D and F.

localization with MY-32 revealed that heterotypic myotubes occurred in hair muscle fibers. This was not surprising since these fibers undergo normal regeneration, in which case a few intertubular did not express neonatal or adult isoforms.

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To further examine the effects of innervation on the expression of embryonic MyHC, injections were analyzed for NCAM and embryonic MyHC. Innervation at 8 weeks after injection into the tibialis anterior muscle [Fig. 3] (immuno-detective localization of 47A and the NCAM-specific polyclonal antibody revealed homotypic muscle fibers that were positive for NCAM but not 47A, or negative for NCAM and positive for 47A]. Therefore, at 8 weeks after myoblast injection, there was still persistence of the embryonic MyHC isoform, even after innervation had occurred. The presence of homotypic muscle fibers that no longer stained for 47A but still stained embryonic MyHC indicates that two developmental pathways of embryonic MyHC can precede innervation. Since there was no correlation between NCAM and embryonic MyHC expression, one can conclude that the developmental switch in embryonic MyHC expression occurs independent of innervation and electrical activity.

To determine if more mature forms of MyHC coexisted with the onset of innervation, homotypic fibers were analyzed with Mabs My-32 (which recognizes all fast MyHC) and BP35 (which recognizes all MyHC isoforms except IIX and embryonic MyHC) in conjunction with NCAM expression [Fig. 4]. Similar to the *in vitro* phenotype of NCAM expression, homotypic fibers reacted with My-32 (but not BP-35), indicating the presence of the IX MyHC isoform. Interestingly, NCAM was colocalized in several myofibers, indicating that innervation had still not occurred. This suggests that, like embryonic MyHC, the expression of the fast IX MyHC isoform is not regulated by innervation. In addition, some homotypic fibers which did become innervated still exhibited a IXK phenotype, suggesting that the expression of other adult MyHCs (such as type I, IIA, or IIB) was not induced by innervation. Combined, these results indicate that the pattern of MyHC expression in L6/AC-14-derived homotypic fibers is not dependent upon innervation and that the development of the mouse muscle fiber phenotype may be governed by an internal control mechanism.

#### Expression of Embryonic MyHC in Heterotypic Fibers

Many of the injection sites also contained muscle fibers which exhibited varying intensities of X-gal labelling, peripherally located nuclei, and bidirectional-shaped cross-sectional areas characteristic of mouse muscle. In addition,  $\beta$ -gal expression was not evenly distributed along the length of these fibers, since areas several hundred micrometers away exhibited little or no staining. These fibers were lo-

cated within the limits of a muscle fascicle, separated from adjacent fibers by a small amount of connective tissue—the endomysium. Based on these criteria these fibers were judged to be the result of donor myoblast fusion to host myoblasts and/or muscle fibers.

To determine if the *in vitro* phenotype of L6 myoblasts was maintained when donor and host nuclei were present to a common cytoplasm, these heterotypic fibers were first examined for the expression of embryonic MyHC—the predominant MyHC isoform expressed by L6 cells in culture (Wiemer et al., 1985; Pia and Mansfield, 1997) and in humomorphic muscle fibers *in vivo*. Localization of 40 labeled cells (20 muscle fibers) at 2 weeks after myoblast transplantation revealed putative heterotypic fibers up to several hundred micrometers away from the injection site [Fig. 5]. Characterization of these fibers using ABC-fluorescent localization with MyHC-specific antibodies demonstrated regionalized expression of embryonic MyHC. These nuclear domains were concentrated around individual nuclei in one area of the fiber and lightly distributed throughout the rest of the cross-sectional area of the fiber. These fibers appeared to be mature since they were not labeled by NMG, which specifically recognizes the isoform of MyHC characteristic of regenerating fibers. Humomorphic fibers in the same area did not express neuronal MyHC, since they did not react with NMG.

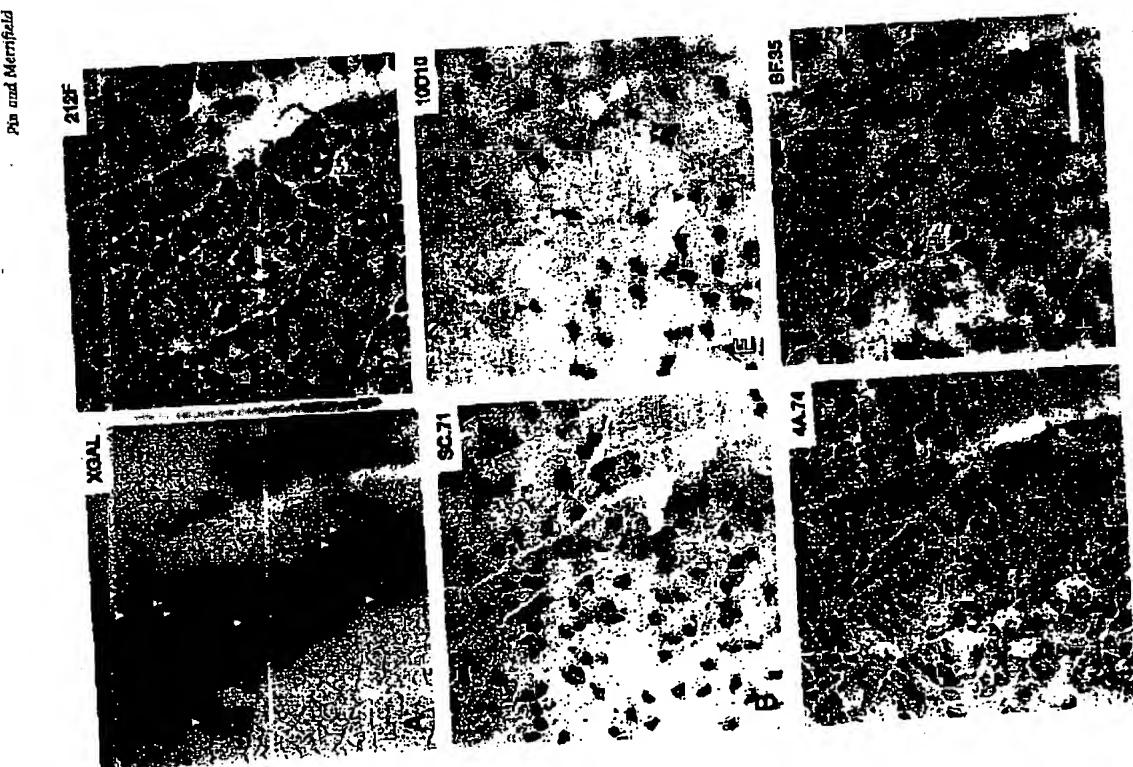
To determine if the expression of embryonic MyHC was transient, injection sites in the tibialis anterior muscle were analyzed at both 6 and 8 weeks after myoblast transplantation. At both 6 and 8 weeks after myoblast transplantation, injection sites in the tibialis anterior muscle showed the persistence of embryonic MyHC. In heterotypic fibers at these later time points similar analyses on the contralateral limb failed to detect embryonic MyHC (data not shown). Interestingly, these nuclear domains were only observed in My-32-positive fibers, suggesting that the regionalized expression of embryonic MyHC may be restricted to fast fiber types. To determine the approximate size of these nuclear domains, longitudinal sections from the tibialis anterior 3 weeks after injection were characterized. When the length of the nuclear domains was determined by measuring the boundaries of the nucleus starting they typically extended 20–25  $\mu\text{m}$  in either direction of an individual nucleus. These results suggest that the embryonic MyHC continues to be expressed for up to 56 days postinjection following the incorporation of L6 myoblasts into fast muscle fibers.

**Expression of the IX MyHC Isoform in Heterotypic Fast Muscle Fibers**

Although the embryonic MyHC isoform was observed in putative heterotypic fibers throughout the course of the

FIG. 7. Characterization of heterotypic fibers 42 days after injection of L6 myoblasts into the tibialis anterior of adult Wistar Retch rats. Serial sections were either stained for X-gal (A) or analyzed with ABC-X-gal immunolocalizations for MyHC- $\beta$ -galactosidase (B, C, D, E, and F). Panel A shows 40/125, C, E, and 10/125, D, fast (IX, IIA, IIB), large area of heterotypic, X-gal-positive fibers. A large area of the muscle fiber is seen to be negative for X-gal. Panel B shows 10/125, C, and 10/125, D, fast (IX, IIA, IIB) and slow (IIC, IIC-I), large area of heterotypic, X-gal-negative fibers. A large area of the muscle fiber is seen to be negative for X-gal. Panel C shows 10/125, C, fast (IX, IIA, IIB) and slow (IIC, IIC-I), large area of heterotypic, X-gal-negative fibers. A large area of the muscle fiber is seen to be negative for X-gal. Panel D shows 10/125, C, fast (IX, IIA, IIB) and slow (IIC, IIC-I), large area of heterotypic, X-gal-positive fibers. A large area of the muscle fiber is seen to be negative for X-gal. Panel E shows 10/125, C, fast (IX, IIA, IIB) and slow (IIC, IIC-I), large area of heterotypic, X-gal-positive fibers. A large area of the muscle fiber is seen to be negative for X-gal. Panel F shows 10/125, C, fast (IX, IIA, IIB) and slow (IIC, IIC-I), large area of heterotypic, X-gal-positive fibers. A large area of the muscle fiber is seen to be negative for X-gal.





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35 or 58 days but not both, similar to fibers in the contralateral limb. No labeling was observed when the injection site was characterized with 1A, 7A, MyHC, or 47A (data not shown), indicating that the fast isoform was not IIA or neonatal MyHC and that embryonic MyHC was not present at this time. Characterization of the injection site 100 days in either direction failed to detect the fast isoform, suggesting that this MyHC was localized to a specific region within the fiber, presumably where fusion of the L6AG-A4 myoblasts had occurred.

To examine whether the expansion of the IX MyHC isoform was maintained in heterotypic fibers transplanted into MyHC, putative heterotypic fibers were examined at 8 weeks after injection. In order to get a large sample size, 8 injection sites within the plantaris muscle were examined. This analysis shows a mixture of different fiber types; this indicates this muscle contains a mixture of different fiber types. Analysis with X-gal immunohistochemistry, revealed a large area of heterotypic fibers (Fig. 9), which were subsequently characterized using AGC-A2 immunohistochemistry with MyHC-specific Mabs. Unfortunately, the percentage of L6/IIA fibers could not be determined since the fibers expressing MyHC did not react with the Mab expressing IX MyHC. However, it was still possible to calculate the proportion of type I, IIA, DA/IX, and IX fibers (Table 3). Even though the percentage of exclusively IX fibers did not show a large increase within the injection site, 19.2% of the fibers expressed both IX and II A MyHC. The total number of fibers which expressed IX MyHC alone or in combination with IIA was 77.6%, an increase of 25.7% over the area outside the injection site.

Upon examination of the injection site with 212F and 10D10, heterotypic fibers coexpressing slow and fast MyHC were not detected. These findings were confirmed by similar observations in the soleus and red gastrocnemius muscles at 8 weeks after injection (data not shown). In addition, slow muscle fibers analyzed over several hundred micrometers using serial sections revealed no heterotypic fast IX MyHC accumulations. Although it is possible that parative nuclear domains of IX or embryonic MyHC could have been overlooked, it seems likely that the fusion of L6 myoblasts to slow fibers resulted in the down-regulation of IX MyHC by L6 muscle.

## DISCUSSION

The introduction of L6AG-A4 myoblasts into a regenerating muscle environment allows these myoblasts to fuse

with each other or with host satellite cells and regenerate muscle fibers to form both heterotypic and homotypic muscle fibers. Homotypic fibers are usually formed by the population of cells which remain at the periphery or in the muscle or between muscle fascicles. Since these myoblasts maintain the characteristic IX MyHC expression, these results and down-regulate embryonic MyHC expression, these results support the hypothesis that L6 myoblasts display a unique potential of forming exclusively fast IX myoblasts both *in vitro* and *in vivo*. The observation that myoblasts maintain their characteristic *in vitro* MyHC profile after maturation *in vivo*, regenerating muscle tissue, was recently done in an avian model [DiMarchi et al., 1992; DiMarchi and Stockdale, 1995], in which primary quail myoblasts of either fast or slow fiber lineage were injected into fetal chick muscle. These injections resulted in the formation of homotypic myotubes which expressed either fast or fast/slow MyHCs in all muscle environments examined. However, these myoblasts were only followed 10 days *in vivo*, since these experiments did not address the possibility of any long-term effects of the environment in general or heterotypic fibers in particular. The observation that L6AG-A2-derived homotypic fibers become intermixed *in vivo* and that this intermixing does not effect the final phenotype of the myotubes, therefore extends the observations previously made in birds. In addition, this is the first demonstration of a fiber-type-specific myoblast cell lineage in mammals.

The intermixing of these fibers is not surprising considering that experiments have been shown to express increasing differentiated myoblasts have been shown to express increased levels of neurotrophic factors [Oppenheim et al., 1993] and higher levels of NCAM [Corradi and Simeoni, 1995]. Two factors known to play a role in muscle fiber intermixing are the interaction [Landmesser et al., 1988] and observation that these homotypic myotubes become intermixed is consistent with experiments that were carried out by Wennig et al. [1991], in which nutritive homotypic fibers apparently became intermixed starting at 4 weeks after injection of cloned neuronal mouse myoblasts into regenerating tissue muscle. Although this group noticed that a fast fiber muscle tissue. Although this group noticed that a fast fiber phenotype predominated early after injection, they also observed a transition to Type I fibers, suggesting that environmental influences may eventually control the phenotype of the myoblasts. These experiments were limited, however, by the fact that the myoblasts used were not characterized *in vitro*, and that the assays used (solid ATPase) were not sensitive enough to delineate between various subtypes of fast fibers. Although we observed intermixed homotypic fast fibers, we did not observe intermixed heterotypic fast fibers. This may be due to the fact that the graft (1 g) contains less than half of the fibers.

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The nature of the mechanisms involved could not be determined by experiments with *Ae. gennadensis*. Since selective transmission by *Leucophenga* was not observed, this approach did not allow us to address the effects of different types of natural enemies on *Leucophenga* parasitism by *Lc* cells.

Hughes and Sian observed that the infected myoblasts adapted in all cases and that the change was based strictly on the environmental cues in which they were placed. However both Cx3 myoblasts (McIndoe and Merrell, 1997) and cultured satellite cells (Jensen et al., 1994) can express a multipotential myoblast stem cell population. Therefore, these donor cell populations appear to become susceptible to environmental cues than the L6 cell line, which has a limited phenotypic potential in culture.

It is possible that MyHC isoforms can form myofibrils, thus allowing the initiation of rhythmic and/or TTX-resistant contraction. The initiation of rhythmic and/or TTX-resistant contraction by donor muscle incorporated into a rat muscle fiber has been reported previously (DeAngelis et al., 1992; Campione et al., 1993). The expression of specifically fast TTX-MyHC isoforms has not been observed, suggesting that some type of restriction may be placed upon slow muscle fibers which prevents the accumulation of TTX-MyHC. One potential mechanism that would allow for the internal regulation of TTX-MyHC involves changes in the expression of the troponin regulatory factors (lmlf). It has been postulated that differential expression of the mlns may be involved in the establishment of slow and fast fiber types with higher amounts of MyoMII cating in fast fibers (Hughes et al., 1993). Indeed, differences in mln expression have been identified in distinct populations of myoblasts, which may relate to their developmental potential *in vivo* (Saitoh et al., 1993; Cuello-DeAngelis et al., 1992; Rudnicki et al., 1993; Cuello-DeAngelis et al., 1993). MyoMII expression is also known to increase MyoMII expression by (de)regulating the synthesis of MyoMII (Saitoh et al., 1993).

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## INTRODUCTION

Vertebrate skeletal muscles are composed of muscle fibers formed from muscle precursor cells, called myoblasts. Within each muscle fiber, a number of muscle-specific contractile proteins are synthesized from families of genes coding multiple protein isoforms. The combinations of genes expressed in particular muscle fibers are large and diverse (Slater and Pett, 1987) giving rise to a multiplicity of muscle fiber phenotypes with unique repertoires of muscle-specific proteins and associated physiological characteristics. In particular, expression of members of the myosin heavy chain (MyHC) multigene family defines fiber type identity and significantly affects fiber contractile properties via myosin ATPase activity (Miesner et al., 1984a,b).

In broad terms, fibers are classified as fast, fast/slow mixed, or slow depending on the presence of MyHC isoforms with fast and/or slow ATPase activities. A central issue in myogenesis is what determines the formation of different muscle fiber types (Stockdale, 1997).

Members of both fast and slow MyHC isoforms occur in nearly all avian skeletal muscle fibers. These isoforms typically involve successive expression of catabolism and then removal and, finally, adult fast MyHC genes (Whalen et al., 1991; Crow and Stockdale, 1986a; Bandmann and Bennett, 1986). However, the slow fast MyHC genes expressed in adult muscle

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